



## $\Box$ 1: XM\_044533. Reports ...[gi:22057705] The record has been replaced by XM\_044533.8

Comment Features Sequence

LOCUS XM\_044533 3766 bp mRNA linear PRI 01-AUG-2002 DEFINITION Homo sapiens sema domain, immunoglobulin domain (Ig), transmembrane

domain (TM) and short cytoplasmic domain, (semaphorin) 4B (SEMA4B), mRNA.

ACCESSION XM\_044533

VERSION XM\_044533.7 GI:22057705

KEYWORDS SOURCE

Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (bases 1 to 3766)

AUTHORS NCBI Annotation Project.

TITLE Direct Submission

JOURNAL Submitted (31-JUL-2002) National Center for Biotechnology

Information, NIH, Bethesda, MD 20894, USA

COMMENT MODEL REFSEQ: This record is predicted by automated computational analysis. This record is derived from a genomic sequence

(NT\_033276) annotated using gene prediction method: BLAST,

supported by mRNA and EST evidence.

Also see:

Documentation of NCBI's Annotation Process

[WARNING] On Jan 5, 2003 this sequence was replaced by qi:27499898.

On Aug 1, 2002 this sequence version replaced gi:20552012.

FEATURES Location/Qualifiers

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/organism="Homo sapiens"
/mol\_type="mRNA"

/db\_xref="taxon:9606" /chromosome="15"

gene 1..3766

/gene="SEMA4B"

/gene\_synonym="KIAA1745; SEMAC; SEMC"

/db\_xref="GeneID:10509"

DS 238..2751

/gene="SEMA4B"

/gene\_synonym="KIAA1745; SEMAC; SEMC"

/codon start=1

/product="similar to Semaphorin 4B"

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misc feature

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/gene="SEMA4B"

/gene\_synonym="KIAA1745; SEMAC; SEMC"

/note="Region: smart00630, Sema, semaphorin domain;

smart00630, Sema, semaphorin domain;"

445..1749 misc\_feature

/gene="SEMA4B"

/gene\_svnonvm="KIAA1745; SEMAC; SEMC" /note="Region: pfam01403, Sema, Sema domain. The Sema

domain occurs in semaphorins, which are a large family of secreted and transmembrane proteins, some of which function as repellent signals during axon guidance. Sema

domains also occur in the hepatocyte growth factor receptor"

misc\_feature

1810..1956

/gene="SEMA4B"

/gene\_synonym="KIAA1745; SEMAC; SEMC"

/note="Region: smart00423, PSI, domain found in Plexins, Semaphorins and Integrins; smart00423, PSI, domain found

in Plexins, Semaphorins and Integrins;"

1810..1881 misc\_feature

/gene="SEMA4B"

/gene\_synonym="KIAA1745; SEMAC; SEMC"

/note="Region: pfam01437, PSI, Plexin repeat. A cysteine rich repeat found in several different extracellular receptors. The function of the repeat is unknown. Three copies of the repeat are found Plexin. Two copies of the repeat are found in mahogany protein. A related C. elegans protein contains four copies of the repeat. The Met receptor contains a single copy of the repeat. The Pfam alignment shows 6 conserved cysteine residues that may form three conserved disulphide bridges, whereas shows 8

conserved cysteines. The pattern of conservation suggests that cysteines 5 and 7 (that are not absolutely conserved) form a disulphide bridge (Personal observation. A

Bateman)"

## ORIGIN

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